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LEUCOCYTE FactsBook ANTIGEN

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Section II THE LEUCOCYTE ANTIGENS

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Molecular weights Polypeptide 48400

unreduced SDS PAGE reduced

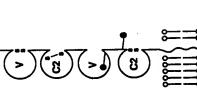
55 kD

55 kD N-linked sites Carbohydrate

Human gene location and size 12pter-p12, 33 kb 1

ם

O-linked



CEV WOC F. CRS WTC

Tissue distribution

2005.90月

CD4 is expressed on most thymocytes and approximately two thirds of human and rat but not in mouse, CD4 is expressed on monocytes and peripheral blood T cells, which constitute the CD8 negative cells 2. macrophages 2.

Structure

crystallography, confirming that they are lg-like 3.4. Domain 2 is characterized by an unusual disulphide within one beta sheet and domain 3 strand C position in domain 2 is replaced with a Trp and there is an extra Cys in the ß strand F 5. The position of the NH2-terminus has been The extracellular domain is made up of four IgSF domains. The structures the amino terminal two domains have been determined by X-ray lacks a disulphide in the position conserved in most IgSF domains. Cat CD4 shows some unusual features with 17 residues inserted between domains 1 and 2.5. There is an additional Cys in domain 1 and the Cys in the unusual $oldsymbol{\beta}$ established for the rat homologue 6. ₹



Function

CD4 is an accessory molecule in the recognition of foreign antigens in association with MHC Class II antigens by T cells ². MAbs against CD4

CD4



motif shown below 8. CD4 is a receptor for HIV-1 (AIDS virus) and the inhibit T cell functions in vivo and in vitro 2. The cytoplasmic domain of CD4 is phosphorylated at Ser residues 408, 415, 431 (see below) when T cells are activated by antigen or phorbol esters 7. The cytoplasmic domain interacts with a lymphocyte-specific tyrosine kinase called p561ck through a binding of the viral gp120 protein is to a region of the amino terminal domain 3,4.



p56lck recognition site (underlined) and Ser residues phosphorylated (in bold) Motifs involved in CD4 function protein (see page 342).



RRDAERMSOI KRLLSE<u>KKTC OC</u>PHRFOKTC SPI (433)

SWISSPROT EMBL/GENBANK REFERENCE M12807 M15768 M13816 P01730 P05540 P06332 AO2109 A02110 A27449 Human Mouse



Amino acid sequence of human CD4

KKVVLGKKGD TVELTCTASQ KKSIQFHWKN SNQIKILGNQ GSFLTKGPSK AFTVEKLTGS GELWWQAERA SSSKSWITFD LKNKEVSVKR VTQDPKLQMG KKLPLHLTLP QALPQYAGSG NITTALEAKT GKLHQEVNLV VMRATQLQKN GOVLLESNIK VLPTWSTPVO PMALIVLGGV AGLLLFIGLG IFFCYRCRHR FANSDTHILD GOSLTITLES PPGSSPSVOC RSPRGKNIDG GKTLSVSQLE LTCEVMGPTS PKLMLSLKLE NKEAKVSKRE KAVHVLNPEA GMWQCLLSDS LNDRADSRRS LWDGGNFPLI IKNLKIEDSD TYICEVEDOK EEVOLLVFGL LODSGTWTCT VLONOKKVEF KIDIVVLAFO KASSIVYKKE GEOVEFSFPL RRQAERMSQI KRLLSEKKTC QCPHRFQKTC SPI MNRGVPFRHL LLVLQLALLP AATOG



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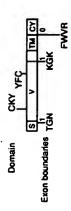
Turner, J.M. et al. (1990) Cell 60, 755-757.

23085 Molecular weights Polypeptide

4 ₹ 90 KD unreduced SDS PAGE reduced

unknown N-linked sites 5 Carbohydrate O-linked

Human gene location and size 2q33-q34, 36 kb 1



Tissue distribution

CD28 is expressed on most T lineage cells and plasma cells 2. Mature thymocytes have higher levels of CD28 than the immature cells and among peripheral T cells, 95% of CD4+ cells and 50% of CD8+ cells are positive 2. Activation of T cells leads to enhanced CD28 expression 2.

Structure

acid level 4. CD28 is particularly similar to CTLA-4 with which it shares a CD28 is a member of the IgSF and is expressed as a disulphide-linked homodimer 23. Human and mouse CD28 are 68% identical at the amino ligand and probably a common ancestor in evolution 4.

Punction

suggesting an important role for CD28 in the interaction between T and B cells. Activation of T cells via CD28 has provided evidence for a CD28 The ligand for CD28 is B7 5.6 which is expressed on activated B cells, signalling pathway which involves stabilization of cytokine mRNA levels and is separate from that used by the TcR-CD3 complex 27.

Database accession numbers

REFERENCE	9	•	90
SWISSPROT EMBL/GENBANK	102988	M34563	X55288
SWISSPROT	P10747		
PIR			
	Human	Mouse	Rat

Amino acid sequence of human CD28

-1 50 100 150 200 202 NKILVKOSPM LVAYDNAVNI SCKYSYNLFS REFRASLHKG LOSAVEVCVV YGNYSQQLQV YSKTGFNCDG KLGNESVTFY LONLYVNQTD IYFCKIEVMY PPPYLONEKS NGTIIHVKGK HLCPSPLFPG PSKPENVLVV VGGYLACYSL LVIVAFIIFW YRSKRSRILH SDYMNMIPRR PGPTRKHYOP YAPPRDFAAY MLRILLIALNE FPSIOVTG

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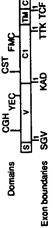
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unknown 60 kD 60 kD Polypeptides 30048 Molecular weights N-linked sites 8 unreduced Carbohydrate SDS PAGE reduced O-linked

сан уес 3q13.3-3q21, 32 kb 1



Tissue distribution

Present on a subset of B cells in vivo and the majority of B cells activated in vitro. Red blood cells, granulocytes, monocytes, resting or activated T cells, thymocytes and platelets do not express B7 2. The antigen is expressed by HTLV-I transformed T cells 3.

glycosylated 4. The sequence of the transmembrane domain is unusual in containing 3 cysteine residues that might be covalently modified or this. The cytoplasmic domain has a preponderance (9/19) of arginine The extracellular domain contains two IgSF domains which are highly participate in intermolecular interactions 4 although there is no evidence for and contains a potential site for calmodulin-dependent phosphorylation (RRES) 4. residues

Function

B7 is the ligand for the CD28 5 and CTLA-4 6 glycoproteins. Cells signals to human T cells, suggesting that the CD28 binding site is upregulated on B cells activated with a variety of agents, including the Epstein-Barr virus 2, cross-linking anti-IgM 2, anti-CD45 and anti-MHC Class II mAbs 9, IL2 and IL4 10. MAbs to B7 block the differentiation of B transfected with either human 7 or murine 8 B7 genes supply co-stimulatory conserved 8. The antigen is not expressed on resting B cells but is strongly cells into antibody secreting cells 11 and the alloactivation of T cells 9 .

1)_[

Comments

This antigen is not related to a mouse antigen called B7 and to avoid confusion the latter is being called B7(2).

Database accession numbers

SWISSPROT EMBL/GENBANK REFERENCE M27533 X60958 PIR Human Mouse

Amino acid sequence of human B7

50 100 150 200 250 262 MSGDMNIWPE YKNRTIFDIT NNLSIVILAL RPSDEGTYEC VVLKYEKDAF ENGEELNAIN TTVSQOPETE-LYAVSSKLOF NMTINHSFMC LIKYGHLRVN OTFNWNTTKQ EHFPON<u>LLPS WAITLISYNG IFVICCLTYC FAP</u>RCRERRR GLSHFCSGVI HVTKEVKEVA TLSCGHNVSV EELADTRIYW QKEKKMVLTM KREHLAEVTL SVKADFPTPS ISDFEIPTSN IRRIICSTSG GFPEPHLSML MGHTRRGGTS PSKCPYLNFF OLLVLA **NERLRRESVR PV**

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